SEQUENCE LISTING

<110> Burnham, Martin K. R.

<120> dexB

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<151> 1997-09-02

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<213> Streptococcus pneumoniae

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Lys Ser Phe Met Asp Ser Asn Gly Asp Gly Val Gly Asp Leu Pro Gly 25

att acc agt aag ttg gac tat cta gct aag tta gga atc aca tcg att

Ile Thr Ser Lys Leu Asp Tyr Leu Ala Lys Leu Gly Ile Thr Ser Ile

35 40 45

tgg ctt tct ccc gtt tat gac agc cct atg gat gat aat ggc tac gat 192

Trp Leu Ser Pro Val Tyr Asp Ser Pro Met Asp Asp Asn Gly Tyr Asp 50 55 60

att gct gat tat caa gcg att gcg gct att ttt gga acc atg gag gac 240

Ile Ala Asp Tyr Gln Ala Ile Ala Ala Ile Phe Gly Thr Met Glu Asp
65 70 75 80

atg gat gaa ctg att gca gaa gct aag aag cgt gat atc cgt atc atc 288

Met Asp Glu Leu Ile Ala Glu Ala Lys Lys Arg Asp Ile Arg Ile Ile 85 90 95

atg gac ttg gtg gtc aat cat acc tcg gat gag cat gcc tgg ttt gta

Met Asp Leu Val Val Asn His Thr Ser Asp Glu His Ala Trp Phe Val 100 105 110

gag gcc tgt gaa aat cct aat agc cct gag cga gac tac tat atc tgg 384

Glu Ala Cys Glu Asn Pro Asn Ser Pro Glu Arg Asp Tyr Tyr Ile Trp
115 120 125

ege gat gaa eee aat gae eta gat tet ate ttt agt ggg tet get tgg 432

Arg Asp Glu Pro Asn Asp Leu Asp Ser Ile Phe Ser Gly Ser Ala Trp
130 135 140

gaa tac gat gaa aag tca ggt caa tac tat etc cae ttt tte age aag 480

Glu Tyr Asp Glu Lys Ser Gly Gln Tyr Tyr Leu His Phe Phe Ser Lys
145 150 155 160

Lys Gln Pro Asp Leu Asn Trp Glu Asn Glu Lys Leu Arg Gln Lys Ile

165 170 175

tat gag atg atg aac ttc tgg att gat aag ggt att ggt ggt ttc cgt
576

Tyr Glu Met Met Asn Phe Trp Ile Asp Lys Gly Ile Gly Gly Phe Arg
180 185 190

atg gat gtt att gac atg att ggc aaa att cct gac gag aag gta gtc

Met Asp Val Ile Asp Met Ile Gly Lys Ile Pro Asp Glu Lys Val Val 195 200 205

aat aat ggt cct atg ctc cat ccc tat ctc aag gaa atg aat cag gcg 672

Asn Asn Gly Pro Met Leu His Pro Tyr Leu Lys Glu Met Asn Gln Ala 210 215 220

acc ttt gga gat aag gat ctc ttg aca gta ggg gag act tgg gga gca 720

Thr Phe Gly Asp Lys Asp Leu Leu Thr Val Gly Glu Thr Trp Gly Ala 225 230 235 240

acg cca gag att gcc aaa ctc tac tct gat cca aag ggg caa gaa ttg 768

Thr Pro Glu Ile Ala Lys Leu Tyr Ser Asp Pro Lys Gly Gln Glu Leu 245 250 255

tot atg gto tto cag ttt gaa cat atc ggt ott cag tat cag gaa ggt 816

Ser Met Val Phe Gln Phe Glu His Ile Gly Leu Gln Tyr Gln Glu Gly 260 265 270

cag cct aaa tgg cac tat caa aaa gag ctg aat atc gct aag tta aaa 864

Gln Pro Lys Trp His Tyr Gln Lys Glu Leu Asn Ile Ala Lys Leu Lys 275 280 285

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Glu Ile Phe Asn Lys Trp Gln Thr Glu Leu Gly Val Glu Asp Gly Trp

1296

290 295 300

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gag	att	ggg	atg	acc	aac	tat	ccg	ttt	gaa	aca	ctg	gat	caa	gta	gaa
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Ala	Arg	Thr	Pro	Met	Gln	Trp	Asp	Glu	Ser	Lys	Asn	Ala	Gly	Phe	Ser
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aca	aat:	caa	cct	taa	$+t\alpha$	aca.	att	aat	cca	aat	tac	gag	ato	atć:	aac

420 425 430

gtc caa gaa gcg ctg gca aat cca gat tct att ttc tat acc tat cag 1344

Val Gln Glu Ala Leu Ala Asn Pro Asp Ser Ile Phe Tyr Thr Tyr Gln 435 440 445

aaa ctg gtc caa att cgc aag gag aat agt tgg cta att cga gct gac 1392

Lys Leu Val Gln Ile Arg Lys Glu Asn Ser Trp Leu Ile Arg Ala Asp 450 460

ttt gaa ttg ctt gat acg gct gat aag gtc ttt gct tat ata cgt aag 1440

Phe Glu Leu Leu Asp Thr Ala Asp Lys Val Phe Ala Tyr Ile Arg Lys 465 470 480

gat ggc gac cgt cgc ttc cta gtt gtg gct aac ttg tcc aat gaa gag 1488

Asp Gly Asp Arg Phe Leu Val Val Ala Asn Leu Ser Asn Glu Glu
485 490 495

caa gac ttg aca gta gaa gga aaa gtc aaa tct gtc ttg att gaa aac 1536

Gln Asp Leu Thr Val Glu Gly Lys Val Lys Ser Val Leu Ile Glu Asn 500 505 510

acc cta gct caa gaa gtc ttt gaa aaa caa atc tta gtt cca tgg gat 1584

Thr Leu Ala Gln Glu Val Phe Glu Lys Gln Ile Leu Val Pro Trp Asp 515 520 525

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1608

Ala Phe Cys Val Glu Leu Leu 530 535

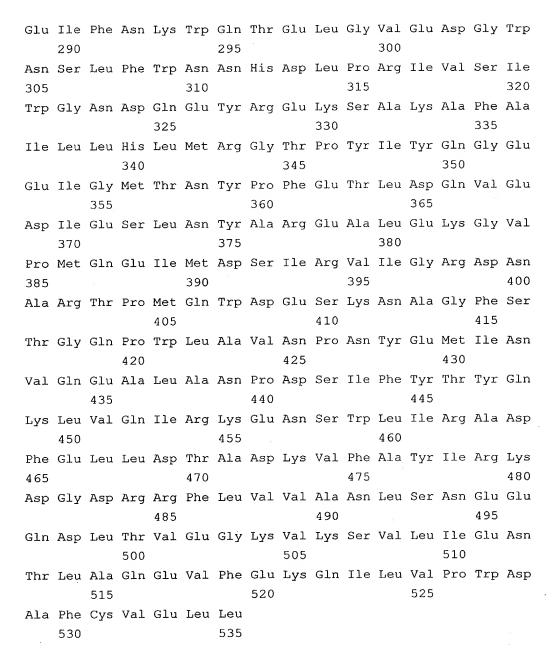
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Ile	Thr	Ser 35	Lys	Leu	Asp	Tyr	Leu 40	Ala	Lys	Leu	Gly	Ile 45	Thr	Ser	Ile
Trp	Leu 50	Ser	Pro	Val	Tyr	Asp 55	Ser	Pro	Met	Asp	Asp 60	Asn	Gly	Tyr	Asp
Ile 65	Ala	Asp	Tyr	Gln	Ala 70	Ile	Ala	Ala	Ile	Phe 75	Gly	Thr	Met	Glu	Asp 80
Met	Asp	Glu	Leu	Ile 85	Ala	Glu	Ala	Lys	Lys 90	Arg	Asp	Ile	Arg	Ile 95	Ile
Met	Asp	Leu	Val 100	Val	Asn	His	Thr	Ser 105	Asp	Glu	His	Ala	Trp 110	Phe	Val
Glu	Ala	Cys 115	Glu	Asn	Pro	Asn	Ser 120	Pro	Glu	Arg	Asp	Tyr 125	Tyr	Ile	Trp
Arg	Asp 130	Glu	Pro	Asn	Asp	Leu 135	Asp	Ser	Ile	Phe	Ser 140	Gly	Ser	Ala	Trp
Glu 145	Tyr	Asp	Glu	Lys	Ser 150	Gly	Gln	Tyr	Tyr	Leu 155	His	Phe	Phe	Ser	Lys 160
Lys	Gln	Pro	Asp	Leu 165	Asn	Trp	Glu	Asn	Glu 170	Lys	Leu	Arg	Gln	Lys 175	Ile
Tyr	Glu	Met	Met 180	Asn	Phe	Trp	Ile	Asp 185	Lys	Gly	Ile	Gly	Gly 190	Phe	Arg
Met _.	Asp	Val 195	Ile	Asp	Met	Ile	Gly 200	Lys	Ile	Pro	Asp ·	Glu 205	Lys	Val	Val
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Thr 225	Phe	Gly	Äsp					Thr			Glu		Trp	Gly	Ala 240
Thr	Pro	Glu	Ile	Ala 245	Lys	Leu	Tyr	Ser	Asp 250	Pro	Lys	Gly	Gln	Glu 255	Leu
Ser	Met	Val	Phe 260	Gln	Phe	Glu	His	Ile 265		Leu	Gln	Tyr	Gln 270	Glu	Gly
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